

=> File .Biotech

=> s (memapsin 2 or beta secretase or beta amyloid precursor protein or APP)
L1 630677 (MEMAPSIN 2 OR BETA SECRETASE OR BETA AMYLOID PRECURSOR PROTEIN
OR APP)

=> s (l1 or memapsin 2) and (OM 99 or OM-99 or dipeptide isostere or dipeptide or isostere) and Alzheimer?

L2 135 (L1 OR MEMAPSIN 2) AND (OM 99 OR OM-99 OR DIPEPTIDE ISOSTERE OR DIPEPTIDE OR ISOSTERE) AND ALZHEIMER?

=> s l2 and (computer program# or software program#)
L3 22 L2 AND (COMPUTER PROGRAM# OR SOFTWARE PROGRAM#)

=> s l3 and (recombinant?)
L4 22 L3 AND (RECOMBINANT?)

=> s l4 and (treat? or therapeut? or diagnos? or prevent?)
5 FILES SEARCHED...
L5 22 L4 AND (TREAT? OR THERAPEUT? OR DIAGNOS? OR PREVENT?)

=> s l5 and (inhibit?)
L6 22 L5 AND (INHIBIT?)

=> s Tang J?/au; s Hong L/au; s Ghosh A/au; s Koelsch G/au
L7 8203 TANG J?/AU

L8 417 HONG L/AU

L9 1933 GHOSH A/AU

L10 100 KOELSCH G/AU

=> s l6 and (l7 or l8 or l9 or l10)
L11 7 L6 AND (L7 OR L8 OR L9 OR L10)

=> d l11 1-7 bib ab

L11 ANSWER 1 OF 7 CAPLUS COPYRIGHT 2003 ACS on STN
AN 2001:12489 CAPLUS
DN 134:80832
TI **Inhibitors of memapsin 2** and use thereof
IN **Tang, Jordan J. N.**; Hong, Ling; Ghosh, Arun K.
PA Oklahoma Medical Research Foundation, USA; The Board of Trustees of the University of Illinois
SO PCT Int. Appl., 86 pp.
CODEN: PIXXD2
DT Patent
LA English
FAN.CNT 3

	PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
PI	WO 2001000665	A2	20010104	WO 2000-US17742	20000627
	WO 2001000665	A3	20010927		
	WO 2001000665	C2	20020725		

W: AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CR, CU, CZ, DE, DK, DM, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, UZ, VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM
RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY,

DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ,
CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG
EP 1194449 A2 20020410 EP 2000-943236 20000627
R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT,
IE, SI, LT, LV, FI, RO

JP 2003506322 T2 20030218 JP 2001-507071 20000627
US 6545127 B1 20030408 US 2000-604608 20000627
US 2002049303 A1 20020425 US 2001-796264 20010228
US 2002164760 A1 20021107 US 2001-795903 20010228
US 2002115600 A1 20020822 US 2001-845226 20010430

PRAI US 1999-141363P P 19990628
US 1999-168060P P 19991130
US 2000-177836P P 20000125
US 2000-178368P P 20000127
US 2000-210292P P 20000608
US 2000-603713 A3 20000627
US 2000-604608 A3 20000627
WO 2000-US17742 W 20000627

AB Methods for the prodn. of purified, catalytically active, **recombinant memapsin 2** have been developed. The substrate and subsite specificity of the catalytically active enzyme have been detd. The substrate and subsite specificity information was used to design substrate analogs of the natural **memapsin 2** substrate that can **inhibit** the function of **memapsin 2**. The substrate analogs are based on peptide sequences, shown to be related to the natural peptide substrates for **memapsin 2**. The substrate analogs contain at least one analog of an amide bond which is not capable of being cleaved by **memapsin 2**. Processes for the synthesis of two substrate analogs including isosteres at the sites of the crit. amino acid residues were developed and the substrate analogs, OMR99-1 and OM99-2, were synthesized. OM99-2 is based on an octapeptide Glu-Val-Asn-Leu-Ala-Ala-Glu-Phe (SEQ ID NO:28) with the Leu-Ala peptide bond substituted by a transition-state **isostere** hydroxyethylene group (Figure 1). The **inhibition** const. of OM99-2 is 1.6×10^{-9} M against **recombinant pro-memapsin 2**. Crystallog. of **memapsin 2** bound to this **inhibitor** was used to det. the three dimensional structure of the protein, as well as the importance of the various residues in binding. This information can be used by those skilled in the art to design new **inhibitors**, using com. available **software programs** and techniques familiar to those in org. chem. and enzymol., to design new **inhibitors** to **memapsin 2**, useful in **diagnostics** and for the **treatment** and/or **prevention** of Alzheimer's disease.

L11 ANSWER 2 OF 7 CAPLUS COPYRIGHT 2003 ACS on STN

AN 2001:12487 CAPLUS

DN 134:68049

TI Catalytically active **recombinant memapsin 2**, 3D crystal structure based **inhibitor** design, synthesis, and screening, for **Alzheimer's disease treatment**

IN **Tang, Jordan J. N.**; Lin, Xinli; Koelsch, Gerald

PA Oklahoma Medical Research Foundation, USA

SO PCT Int. Appl., 87 pp.

CODEN: PIXXD2

DT Patent

LA English

FAN.CNT 3

	PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
PI	WO 2001000663	A2	20010104	WO 2000-US17661	20000627
	WO 2001000663	A3	20011004		

W: AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CR, CU,
CZ, DE, DK, DM, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL,

IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA,
 MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI,
 SK, SL, TJ, TM, TR, TT, TZ, UA, UG, UZ, VN, YU, ZA, ZW, AM, AZ,
 BY, KG, KZ, MD, RU, TJ, TM
 RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY,
 DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ,
 CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG

EP 1196609 A2 20020417 EP 2000-943208 20000627
 R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT,
 IE, SI, LT, LV, FI, RO

JP 2003503072	T2	20030128	JP 2001-507069	20000627
US 6545127	B1	20030408	US 2000-604608	20000627
US 2002049303	A1	20020425	US 2001-796264	20010228
US 2002164760	A1	20021107	US 2001-795903	20010228
US 2002115600	A1	20020822	US 2001-845226	20010430

PRAI US 1999-141363P P 19990628
 US 1999-168060P P 19991130
 US 2000-177836P P 20000125
 US 2000-178368P P 20000127
 US 2000-210292P P 20000608
 US 2000-603713 A3 20000627
 US 2000-604608 A3 20000627
 WO 2000-US17661 W 20000627

AB A method for producing catalytically active **recombinant memapsin 2** comprising expression in a bacteria and refolding the **recombinant memapsin 2** under conditions which dissociate and then slowly refold the enzyme into a catalytically active form is disclosed. A method of isolating **inhibitors** of cleavage by **memapsin 2** comprising adding to one or more potential **inhibitors** of catalytically active **recombinant memapsin 2**, and a substrate for **memapsin 2**, and screening for decreased cleavage of the substrate by the **inhibitors**, wherein the **inhibitors** are in a library of small synthetic molecules, like proteins and peptides. Alternatively, the **inhibitors** are oligonucleotides **preventing** or decreasing expression of catalytically active **memapsin 2**. A method for designing or obtaining **inhibitors** of catalytically active **memapsin 2** comprising modeling an **inhibitor** based on the crystal coordinates of **memapsin 2** or parameters. A database comprising binding properties and chemical structures of compounds designed or screened by modeling an **inhibitor** based on the crystal coordinates of **memapsin 2** or parameters is claimed. A method of **treating** or **preventing** **Alzheimer's** disease comprising administering to a patient in need thereof an **inhibitor** of **memapsin 2** which binds to the active site of the **memapsin 2** defined by the presence of two catalytic aspartic residues and substrate binding cleft, is also claimed. The cDNAs of two new human membrane-associated aspartic proteases, **memapsin 1** and **memapsin 2**, have been cloned and sequenced. The substrate and subsite specificity of the catalytically active enzyme have been determined. The substrate and subsite specificity information was used to design substrate analogs of the natural **memapsin 2** substrate that can **inhibit** the function of **memapsin 2**. The substrate analogs are based on peptide sequences, shown to be related to the natural peptide substrates for **memapsin 2**. The substrate analogs contain at least one analog of an amide bond which is not capable of being cleaved by **memapsin 2**. Processes for the synthesis of two substrate analogs including isosteres at the sites of the critical amino acid residues were developed and the substrate analogs, OM99-1 and OM99-2, were synthesized. OM99-2 is based on an octapeptide Glu-Val-Asn-Leu-Ala-Ala-Glu-Phe (SEQ ID NO:28) with the Leu-Ala peptide bond substituted by a transition-state **isostere** hydroxyethylene group (Fig. 1). The **inhibition** constant of OM99-2 is 1.6×10^9 M

against **recombinant pro-memapsin 2**.
Crystallog. of **memapsin 2** bound to this
inhibitor was used to det. the three dimensional structure of the
protein, as well as the importance of the various residues in binding.
This information can be used to design new **inhibitors**, using
com. available **software programs** and techniques
familiar to those in org. chem. and enzymol., to design new
inhibitors to **memapsin 2**, useful in
diagnostics and for the **treatment** and/or
prevention of **Alzheimer's** disease.

L11 ANSWER 3 OF 7 USPATFULL on STN
AN 2003:134541 USPATFULL
TI **Inhibitors** of **memapsin 2** and use thereof
IN **Tang, Jordan J. N.**, Edmond, OK, UNITED STATES
Koelsch, Gerald, Oklahoma City, OK, UNITED STATES
Ghosh, Arun K., River Forest, IL, UNITED STATES
PA Oklahoma Medical Research Foundation, Oklahoma City, OK (U.S.
corporation)
PI US 2003092629 A1 20030515
AI US 2001-32818 A1 20011228 (10)
PRAI US 2001-275756P 20010314 (60)
US 2000-258705P 20001228 (60)
DT Utility
FS APPLICATION
LREP HAMILTON, BROOK, SMITH & REYNOLDS, P.C., 530 VIRGINIA ROAD, P.O. BOX
9133, CONCORD, MA, 01742-9133
CLMN Number of Claims: 24
ECL Exemplary Claim: 1
DRWN 9 Drawing Page(s)
LN.CNT 2203
CAS INDEXING IS AVAILABLE FOR THIS PATENT.
AB Methods for the production of purified, catalytically active,
recombinant memapsin 2 have been developed-,
The substrate and subsite specificity of the catalytically active enzyme
have been determined by a method which determines the initial hydrolysis
rate of the substrates by using MALDI-TOF/MS. Alternatively, the subsite
specificity of **memapsin** can be determined by probing a library of
inhibitors with **memapsin 2** and subsequently
detecting the bound **memapsin 2** with an antibody
raised to **memapsin 2** and an alkaline phosphatase
conjugated secondary antibody. The substrate and subsite specificity
information was used to design substrate analogs of the natural
memapsin 2 substrate that can **inhibit** the
function of **memapsin 2**. The substrate analogs are
based on peptide sequences, shown to be related to the natural peptide
substrates for **memapsin 2**. The substrate analogs
contain at least one analog of an amide bond which is not capable of
being cleaved by **memapsin 2**. Processes for the
synthesis of substrate analogues including isosteres at the sites of the
critical amino acid residues were developed and the more than seventy
substrate analogues were synthesized, among which MMI-005, MMI-012,
MMI-017, MMI-018, MMI-025, MMI-026, MMI-037, MMI-039, MMI-040, MMI-066,
MMI-070, and MMI-071 have **inhibition** constants in the range of
1.4-61.4.times.10.sup.-9 M against **recombinant pro-**
memapsin 2. These **inhibitors** are useful in
diagnostics and for the **treatment** and/or
prevention of **Alzheimer's** disease.

L11 ANSWER 4 OF 7 USPATFULL on STN
AN 2003:96167 USPATFULL
TI Catalytically active **recombinant memapsin** and methods of use
thereof
IN **Tang, Jordan J. N.**, Edmond, OK, United States
Lin, Xinli, Edmond, OK, United States

Koelsch, Gerald, Oklahoma City, OK, United States
Hong, Lin, Oklahoma City, OK, United States
PA Oklahoma Medical Research Foundation, Oklahoma City, OK, United States
(U.S. corporation)
PI US 6545127 B1 20030408
AI US 2000-604608 20000627 (9)
PRAI US 1999-141363P 19990628 (60)
US 1999-168060P 19991130 (60)
US 2000-177836P 20000125 (60)
US 2000-178368P 20000127 (60)
US 2000-210292P 20000608 (60)
DT Utility
FS GRANTED
EXNAM Primary Examiner: Borin, Michael; Assistant Examiner: Zhou, Shuba
LREP Hamilton, Brook, Smith & Reynolds, P.C.
CLMN Number of Claims: 18
ECL Exemplary Claim: 7
DRWN 21 Drawing Figure(s); 12 Drawing Page(s)
LN.CNT 2563
CAS INDEXING IS AVAILABLE FOR THIS PATENT.
AB Methods for the production of purified, catalytically active, **recombinant memapsin 2** have been developed. The substrate and subsite specificity of the catalytically active enzyme have been determined. The substrate and subsite specificity information was used to design substrate analogs of the natural **memapsin 2** substrate that can **inhibit** the function of **memapsin 2**. The substrate analogs are based on peptide sequences, shown to be related to the natural peptide substrates for **memapsin 2**. The substrate analogs contain at least one analog of an amide bond which is not capable of being cleaved by **memapsin 2**. Processes for the synthesis of two substrate analogues including isosteres at the sites of the critical amino acid residues were developed and the substrate analogues, OMR99-1 and OMR99-2, were synthesized. OMR99-2 is based on an octapeptide Glu-Val-Asn-Leu-Ala-Ala-Glu-Phe (SEQ ID NO:28) with the Leu-Ala peptide bond substituted by a transition-state **isostere** hydroxyethylene group (FIG. 1). The **inhibition** constant of OMR99-2 is 1.6.times.10.sup.-9M against **recombinant pro-memapsin 2**. Crystallography of **memapsin 2** bound to this **inhibitor** was used to determine the three dimensional structure of the protein, as well as the importance of the various residues in binding. This information can be used by those skilled in the art to design new **inhibitors**, using commercially available **software programs** and techniques familiar to those in organic chemistry and enzymology, to design new **inhibitors** to **memapsin 2**, useful in **diagnostics** and for the **treatment** and/or **prevention** of **Alzheimer's** disease.

L11 ANSWER 5 OF 7 USPATFULL on STN
AN 2002:294717 USPATFULL
TI Catalytically active **recombinant memapsin** and methods of use thereof
IN Lin, Xinli, Edmond, OK, UNITED STATES
Koelsch, Gerald, Oklahoma City, OK, UNITED STATES
Tang, Jordan J.N., Edmond, OK, UNITED STATES
PA Oklahoma Medical Research Foundation
PI US 2002164760 A1 20021107
AI US 2001-795903 A1 20010228 (9)
RLI Division of Ser. No. US 2000-604608, filed on 27 Jun 2000, PENDING
PRAI US 1999-141363P 19990628 (60)
US 1999-168060P 19991130 (60)
US 2000-177836P 20000125 (60)
US 2000-178368P 20000127 (60)
US 2000-210292P 20000608 (60)

DT Utility
FS APPLICATION
LREP PATREA L. PABST, HOLLAND & KNIGHT LLP, SUITE 2000, ONE ATLANTIC CENTER,
1201 WEST PEACHTREE STREET, N.E., ATLANTA, GA, 30309-3400
CLMN Number of Claims: 33
ECL Exemplary Claim: 1
DRWN 12 Drawing Page(s)
LN.CNT 2440
CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Methods for the production of purified, catalytically active,
recombinant memapsin 2 have been developed.
The substrate and subsite specificity of the catalytically active enzyme
have been determined. The substrate and subsite specificity information
was used to design substrate analogs of the natural **memapsin**
2 substrate that can **inhibit** the function of
memapsin 2. The substrate analogs are based on peptide
sequences, shown to be related to the natural peptide substrates for
memapsin 2. The substrate analogs contain at least one
analog of an amide bond which is not capable of being cleaved by
memapsin 2. Processes for the synthesis of two
substrate analogues including isosteres at the sites of the critical
amino acid residues were developed and the substrate analogues, OMR99-1
and OM99-2, were synthesized. OM99-2 is based on an octapeptide
Glu-Val-Asn-Leu-Ala-Ala-Glu-Phe (SEQ ID NO:28) with the Leu-Ala peptide
bond substituted by a transition-state **isostere**
hydroxyethylene group (FIG. 1). The **inhibition** constant of
OM99-2 is 1.6.times.10.sup.-9 M against **recombinant pro-**
memapsin 2. Crystallography of **memapsin**
2 bound to this **inhibitor** was used to determine the
three dimensional structure of the protein, as well as the importance of
the various residues in binding. This information can be used by those
skilled in the art to design new **inhibitors**, using
commercially available **software programs** and
techniques familiar to those in organic chemistry and enzymology, to
design new **inhibitors** to **memapsin 2**,
useful in **diagnostics** and for the **treatment** and/or
prevention of Alzheimer's disease.

L11 ANSWER 6 OF 7 USPATFULL on STN
AN 2002:214213 USPATFULL
TI **Inhibitors** of **memapsin 2** and use thereof
IN Koelsch, Gerald, Oklahoma City, OK, UNITED STATES
Tang, Jordan J.N., Edmond, OK, UNITED STATES
Hong, Lin, Oklahoma City, OK, UNITED STATES
Ghosh, Arun K., River Forest, IL, UNITED STATES
PA Oklahoma Medical Research Foundation (U.S. corporation)
PI US 2002115600 A1 20020822
AI US 2001-845226 A1 20010430 (9)
RLI Division of Ser. No. US 2000-603713, filed on 27 Jun 2000, PENDING
PRAI US 1999-141363P 19990628 (60)
US 1999-168060P 19991130 (60)
US 2000-177836P 20000125 (60)
US 2000-178368P 20000127 (60)
US 2000-210292P 20000608 (60)

DT Utility
FS APPLICATION
LREP Patrea L. Pabst, Arnall Golden & Gregory, LLP, 2800 One Atlantic Center,
1201 West Peachtree Street, Atlanta, GA, 30309-3450
CLMN Number of Claims: 23
ECL Exemplary Claim: 1
DRWN 12 Drawing Page(s)
LN.CNT 2377
CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Methods for the production of purified, catalytically active,
recombinant memapsin 2 have been developed.

The substrate and subsite specificity of the catalytically active enzyme have been determined. The substrate and subsite specificity information was used to design substrate analogs of the natural **memapsin 2** substrate that can **inhibit** the function of **memapsin 2**. The substrate analogs are based on peptide sequences, shown to be related to the natural peptide substrates for **memapsin 2**. The substrate analogs contain at least one analog of an amide bond which is not capable of being cleaved by **memapsin 2**. Processes for the synthesis of two substrate analogues including isosteres at the sites of the critical amino acid residues were developed and the substrate analogues, OMR99-1 and OM99-2, were synthesized. OM99-2 is based on an octapeptide Glu-Val-Asn-Leu-Ala-Ala-Glu-Phe (SEQ ID NO:28) with the Leu-Ala peptide bond substituted by a transition-state **isostere** hydroxyethylene group (FIG. 1). The **inhibition** constant of OM99-2 is 1.6.times.10.sup.-9 M against **recombinant pro-memapsin 2**. Crystallography of **memapsin 2** bound to this **inhibitor** was used to determine the three dimensional structure of the protein, as well as the importance of the various residues in binding. This information can be used by those skilled in the art to design new **inhibitors**, using commercially available **software programs** and techniques familiar to those in organic chemistry and enzymology, to design new **inhibitors** to **memapsin 2**, useful in **diagnostics** and for the **treatment** and/or **prevention** of Alzheimer's disease.

L11 ANSWER 7 OF 7 USPATFULL on STN

AN 2002:92777 USPATFULL

TI Catalytically active **recombinant** memapsin and methods of use thereof

IN **Tang, Jordan J. N.**, Edmond, OK, UNITED STATES
 Lin, Xinli, Edmond, OK, UNITED STATES
 Koelsch, Gerald, Oklahoma City, OK, UNITED STATES
 Hong, Lin, Oklahoma City, OK, UNITED STATES

PI US 2002049303 A1 20020425

AI US 2001-796264 A1 20010228 (9)

RLI Division of Ser. No. US 2000-604608, filed on 27 Jun 2000, PENDING

PRAI US 1999-141363P 19990628 (60)

US 1999-168060P 19991130 (60)

US 2000-177836P 20000125 (60)

US 2000-178368P 20000127 (60)

DT Utility

FS APPLICATION

LREP PATREA L. PABST, HOLLAND & KNIGHT LLP, SUITE 2000, ONE ATLANTIC CENTER, 1201 WEST PEACHTREE STREET, N.E., ATLANTA, GA, 30309-3400

CLMN Number of Claims: 33

ECL Exemplary Claim: 1

DRWN 12 Drawing Page(s)

LN.CNT 2441

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Methods for the production of purified, catalytically active, **recombinant memapsin 2** have been developed. The substrate and subsite specificity of the catalytically active enzyme have been determined. The substrate and subsite specificity information was used to design substrate analogs of the natural **memapsin 2** substrate that can **inhibit** the function of **memapsin 2**. The substrate analogs are based on peptide sequences, shown to be related to the natural peptide substrates for **memapsin 2**. The substrate analogs contain at least one analog of an amide bond which is not capable of being cleaved by **memapsin 2**. Processes for the synthesis of two substrate analogs including isosteres at the sites of the critical amino acid residues were developed and the substrate analogs, OMR99-1 and OM99-2, were synthesized. OM99-2 is based on an octapeptide

Glu-Val-Asn-Leu-Ala-Ala-Glu-Phe (SEQ ID NO:28) with the Leu-Ala peptide bond substituted by a transition-state **isostere** hydroxyethylene group (FIG. 1). The **inhibition** constant of OM99-2 is 1.6.times.10.sup.-9 M against **recombinant** pro-**memapsin 2**. Crystallography of **memapsin 2** bound to this **inhibitor** was used to determine the three dimensional structure of the protein, as well as the importance of the various residues in binding. This information can be used by those skilled in the art to design new **inhibitors**, using commercially available **software programs** and techniques familiar to those in organic chemistry and enzymology, to design new **inhibitors** to **memapsin 2**, useful in **diagnostics** and for the **treatment** and/or **prevention** of **Alzheimer's** disease.

=> d his

(FILE 'HOME' ENTERED AT 16:27:26 ON 05 AUG 2003)

FILE 'MEDLINE, CAPLUS, BIOSIS, BIOTECHDS, EMBASE, USPATFULL, WPIDS'
ENTERED AT 16:29:30 ON 05 AUG 2003

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L1      630677 S (MEMAPSIN 2 OR BETA SECRETASE OR BETA AMYLOID PRECURSOR PROTE
L2      135 S (L1 OR MEMAPSIN 2) AND (OM 99 OR OM-99 OR DIPEPTIDE ISOSTERE
L3      22 S L2 AND (COMPUTER PROGRAM# OR SOFTWARE PROGRAM#)
L4      22 S L3 AND (RECOMBINANT?)
L5      22 S L4 AND (TREAT? OR THERAPEUT? OR DIAGNOS? OR PREVENT?)
L6      22 S L5 AND (INHIBIT?)
L7      8203 S TANG J?/AU
L8      417 S HONG L/AU
L9      1933 S GHOSH A/AU
L10     100 S KOELSCH G/AU
L11     7 S L6 AND (L7 OR L8 OR L9 OR L10)
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=> d l6 1-22 bib ab

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L6      ANSWER 1 OF 22  CAPLUS  COPYRIGHT 2003 ACS on STN
AN      2001:12489  CAPLUS
DN      134:80832
TI      Inhibitors of memapsin 2 and use thereof
IN      Tang, Jordan J. N.; Hong, Ling; Ghosh, Arun K.
PA      Oklahoma Medical Research Foundation, USA; The Board of Trustees of the
        University of Illinois
SO      PCT Int. Appl., 86 pp.
        CODEN: PIXXD2
DT      Patent
LA      English
FAN.CNT 3
```

	PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
PI	WO 2001000665	A2	20010104	WO 2000-US17742	20000627
	WO 2001000665	A3	20010927		
	WO 2001000665	C2	20020725		
	W:	AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CR, CU, CZ, DE, DK, DM, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, UZ, VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM			
	RW:	GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG			
EP	1194449	A2	20020410	EP 2000-943236	20000627
	R:	AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT, IE, SI, LT, LV, FI, RO			

	JP 2003506322	T2	20030218	JP 2001-507071	20000627
	US 6545127	B1	20030408	US 2000-604608	20000627
	US 2002049303	A1	20020425	US 2001-796264	20010228
	US 2002164760	A1	20021107	US 2001-795903	20010228
	US 2002115600	A1	20020822	US 2001-845226	20010430
PRAI	US 1999-141363P	P	19990628		
	US 1999-168060P	P	19991130		
	US 2000-177836P	P	20000125		
	US 2000-178368P	P	20000127		
	US 2000-210292P	P	20000608		
	US 2000-603713	A3	20000627		
	US 2000-604608	A3	20000627		
	WO 2000-US17742	W	20000627		
AB	<p>Methods for the prodn. of purified, catalytically active, recombinant memapsin 2 have been developed. The substrate and subsite specificity of the catalytically active enzyme have been detd. The substrate and subsite specificity information was used to design substrate analogs of the natural memapsin 2 substrate that can inhibit the function of memapsin 2. The substrate analogs are based on peptide sequences, shown to be related to the natural peptide substrates for memapsin 2. The substrate analogs contain at least one analog of an amide bond which is not capable of being cleaved by memapsin 2. Processes for the synthesis of two substrate analogs including isosteres at the sites of the crit. amino acid residues were developed and the substrate analogs, OMR99-1 and OMR99-2, were synthesized. OMR99-2 is based on an octapeptide Glu-Val-Asn-Leu-Ala-Ala-Glu-Phe (SEQ ID NO:28) with the Leu-Ala peptide bond substituted by a transition-state isostere hydroxyethylene group (Figure 1). The inhibition const. of OMR99-2 is 1.6×10^{-9} M against recombinant pro-memapsin 2. Crystallog. of memapsin 2 bound to this inhibitor was used to det. the three dimensional structure of the protein, as well as the importance of the various residues in binding. This information can be used by those skilled in the art to design new inhibitors, using com. available software programs and techniques familiar to those in org. chem. and enzymol., to design new inhibitors to memapsin 2, useful in diagnostics and for the treatment and/or prevention of Alzheimer's disease.</p>				

L6 ANSWER 2 OF 22 CAPLUS COPYRIGHT 2003 ACS on STN
 AN 2001:12487 CAPLUS
 DN 134:68049

TI Catalytically active **recombinant memapsin 2**,
 3D crystal structure based **inhibitor** design, synthesis, and
 screening, for **Alzheimer's** disease **treatment**

IN Tang, Jordan J. N.; Lin, Xinli; Koelsch, Gerald

PA Oklahoma Medical Research Foundation, USA

SO PCT Int. Appl., 87 pp.

CODEN: PIXXD2

DT Patent

LA English

FAN.CNT 3

	PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
PI	WO 2001000663	A2	20010104	WO 2000-US17661	20000627
	WO 2001000663	A3	20011004		
	W: AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CR, CU, CZ, DE, DK, DM, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, UZ, VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY,				

DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ,
 CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG
 EP 1196609 A2 20020417 EP 2000-943208 20000627
 R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT,
 IE, SI, LT, LV, FI, RO

JP 2003503072	T2	20030128	JP 2001-507069	20000627
US 6545127	B1	20030408	US 2000-604608	20000627
US 2002049303	A1	20020425	US 2001-796264	20010228
US 2002164760	A1	20021107	US 2001-795903	20010228
US 2002115600	A1	20020822	US 2001-845226	20010430

PRAI US 1999-141363P P 19990628
 US 1999-168060P P 19991130
 US 2000-177836P P 20000125
 US 2000-178368P P 20000127
 US 2000-210292P P 20000608
 US 2000-603713 A3 20000627
 US 2000-604608 A3 20000627
 WO 2000-US17661 W 20000627

AB A method for producing catalytically active **recombinant memapsin 2** comprising expression in a bacteria and refolding the **recombinant memapsin 2** under conditions which dissociate and then slowly refold the enzyme into a catalytically active form is disclosed. A method of isolating **inhibitors** of cleavage by **memapsin 2** comprising adding to one or more potential **inhibitors** of catalytically active **recombinant memapsin 2**, and a substrate for **memapsin 2**, and screening for decreased cleavage of the substrate by the **inhibitors**, wherein the **inhibitors** are in a library of small synthetic molecules, like proteins and peptides. Alternatively, the **inhibitors** are oligonucleotides preventing or decreasing expression of catalytically active **memapsin 2**. A method for designing or obtaining **inhibitors** of catalytically active **memapsin 2** comprising modeling an **inhibitor** based on the crystal coordinates of **memapsin 2** or parameters. A database comprising binding properties and chemical structures of compounds designed or screened by modeling an **inhibitor** based on the crystal coordinates of **memapsin 2** or parameters is claimed. A method of treating or preventing Alzheimer's disease comprising administering to a patient in need thereof an **inhibitor** of **memapsin 2** which binds to the active site of the **memapsin 2** defined by the presence of two catalytic aspartic residues and substrate binding cleft, is also claimed. The cDNAs of two new human membrane-associated aspartic proteases, **memapsin 1** and **memapsin 2**, have been cloned and sequenced. The substrate and subsite specificity of the catalytically active enzyme have been determined. The substrate and subsite specificity information was used to design substrate analogs of the natural **memapsin 2** substrate that can inhibit the function of **memapsin 2**. The substrate analogs are based on peptide sequences, shown to be related to the natural peptide substrates for **memapsin 2**. The substrate analogs contain at least one analog of an amide bond which is not capable of being cleaved by **memapsin 2**. Processes for the synthesis of two substrate analogs including isosteres at the sites of the critical amino acid residues were developed and the substrate analogs, OMR99-1 and OMR99-2, were synthesized. OMR99-2 is based on an octapeptide Glu-Val-Asn-Leu-Ala-Ala-Glu-Phe (SEQ ID NO:28) with the Leu-Ala peptide bond substituted by a transition-state isostere hydroxyethylene group (Fig. 1). The inhibition constant of OMR99-2 is 1.6×10^9 M against recombinant pro-**memapsin 2**. Crystallography of **memapsin 2** used to this **inhibitor** was used to determine the three dimensional structure of the protein, as well as the importance of the various residues in binding. This information can be used to design new **inhibitors**, using

com. available **software programs** and techniques familiar to those in org. chem. and enzymol., to design new **inhibitors to memapsin 2**, useful in **diagnostics** and for the **treatment** and/or **prevention of Alzheimer's disease**.

L6 ANSWER 3 OF 22 USPATFULL on STN
AN 2003:200784 USPATFULL
TI Immunogenic HBc chimer particles having enhanced stability
IN Birkett, Ashley J., Escondido, CA, UNITED STATES
PI US 2003138769 A1 20030724
AI US 2001-930915 A1 20010815 (9)
RLI Continuation-in-part of Ser. No. US 2000-226867, filed on 22 Aug 2000, PENDING Continuation-in-part of Ser. No. US 2000-225843, filed on 16 Aug 2000, PENDING
DT Utility
FS APPLICATION
LREP WELSH & KATZ, LTD, 120 S RIVERSIDE PLAZA, 22ND FLOOR, CHICAGO, IL, 60606
CLMN Number of Claims: 115
ECL Exemplary Claim: 1
DRWN 10 Drawing Page(s)
LN.CNT 6993
AB A chimeric, carboxy-terminal truncated hepatitis B virus nucleocapsid protein (HBc) is disclosed that is engineered for both enhanced stability of self-assembled particles and the display of an immunogenic epitope. The display of the immunogenic epitope is displayed in the immunogenic loop of HBc, whereas the enhanced stability of self-assembled particles is obtained by the presence of at least one heterologous cysteine residue near the carboxy-terminus of the chimera molecule. Methods of making and using the chimeras are also disclosed.

L6 ANSWER 4 OF 22 USPATFULL on STN
AN 2003:187895 USPATFULL
TI 12 human secreted proteins
IN Ni, Jian, Germantown, MD, UNITED STATES
Young, Paul E., Gaithersburg, MD, UNITED STATES
Kenny, Joseph J., Damascus, MD, UNITED STATES
Olsen, Henrik S., Gaithersburg, MD, UNITED STATES
Moore, Paul A., Germantown, MD, UNITED STATES
Wei, Ying-Fei, Berkeley, CA, UNITED STATES
Greene, John M., Gaithersburg, MD, UNITED STATES
Ruben, Steven M., Olney, MD, UNITED STATES
PI US 2003129685 A1 20030710
AI US 2001-836353 A1 20010418 (9)
RLI Continuation-in-part of Ser. No. WO 1999-US25031, filed on 27 Oct 1999, UNKNOWN
PRAI US 1998-105971P 19981028 (60)
US 2000-198407P 20000419 (60)
DT Utility
FS APPLICATION
LREP HUMAN GENOME SCIENCES INC, 9410 KEY WEST AVENUE, ROCKVILLE, MD, 20850
CLMN Number of Claims: 23
ECL Exemplary Claim: 1
DRWN 59 Drawing Page(s)
LN.CNT 31945
CAS INDEXING IS AVAILABLE FOR THIS PATENT.
AB The present invention relates to novel human secreted proteins and isolated nucleic acids containing the coding regions of the genes encoding such proteins. Also provided are vectors, host cells, antibodies, and **recombinant** methods for producing human secreted proteins. The invention further relates to **diagnostic** and **therapeutic** methods useful for **diagnosing** and **treating** diseases, disorders, and/or conditions related to these novel human secreted proteins.

L6 ANSWER 5 OF 22 USPATFULL on STN
 AN 2003:165862 USPATFULL
 TI Directed evolution of novel binding proteins
 IN Ladner, Robert Charles, Ijamsville, MD, UNITED STATES
 Guterman, Sonia Kosow, Belmont, MA, UNITED STATES
 Roberts, Bruce Lindsay, Milford, MA, UNITED STATES
 Markland, William, Milford, MA, UNITED STATES
 Ley, Arthur Charles, Newton, MA, UNITED STATES
 Kent, Rachel Baribault, Boxborough, MA, UNITED STATES
 PI US 2003113717 A1 20030619
 AI US 2001-893878 A1 20010629 (9)
 RLI Continuation of Ser. No. US 1997-993776, filed on 18 Dec 1997, PENDING
 Continuation of Ser. No. US 1995-415922, filed on 3 Apr 1995, PATENTED
 Continuation of Ser. No. US 1993-9319, filed on 26 Jan 1993, PATENTED
 Division of Ser. No. US 1991-664989, filed on 1 Mar 1991, PATENTED
 Continuation-in-part of Ser. No. US 1990-487063, filed on 2 Mar 1990,
 ABANDONED Continuation-in-part of Ser. No. US 1988-240160, filed on 2
 Sep 1988, ABANDONED
 PRAI WO 1989-US3731 19890901
 DT Utility
 FS APPLICATION
 LREP BROWDY AND NEIMARK, P.L.L.C., 624 Ninth Street, N.W., Washington, DC,
 20001
 CLMN Number of Claims: 25
 ECL Exemplary Claim: 1
 DRWN 16 Drawing Page(s)
 LN.CNT 15933
 CAS INDEXING IS AVAILABLE FOR THIS PATENT.
 AB In order to obtain a novel binding protein against a chosen target, DNA
 molecules, each encoding a protein comprising one of a family of similar
 potential binding domains and a structural-signal calling for the
 display of the protein on the outer surface of a chosen bacterial cell,
 bacterial spore or phage (genetic package) are introduced into a genetic
 package. The protein is expressed and the potential binding domain is
 displayed on the outer surface of the package. The cells or viruses
 bearing the binding domains which recognize the target molecule are
 isolated and amplified. The successful binding domains are then
 characterized. One or more of these successful binding domains is used
 as a model for the design of a new family of potential binding domains,
 and the process is repeated until a novel binding domain having a
 desired affinity for the target molecule is obtained. In one embodiment,
 the first family of potential binding domains is related to bovine
 pancreatic trypsin **inhibitor**, the genetic package is M13
 phage, and the protein includes the outer surface transport signal of
 the M13 gene III protein.

L6 ANSWER 6 OF 22 USPATFULL on STN
 AN 2003:134541 USPATFULL
 TI **Inhibitors of memapsin 2** and use thereof
 IN Tang, Jordan J. N., Edmond, OK, UNITED STATES
 Koelsch, Gerald, Oklahoma City, OK, UNITED STATES
 Ghosh, Arun K., River Forest, IL, UNITED STATES
 PA Oklahoma Medical Research Foundation, Oklahoma City, OK (U.S.
 corporation)
 PI US 2003092629 A1 20030515
 AI US 2001-32818 A1 20011228 (10)
 PRAI US 2001-275756P 20010314 (60)
 US 2000-258705P 20001228 (60)
 DT Utility
 FS APPLICATION
 LREP HAMILTON, BROOK, SMITH & REYNOLDS, P.C., 530 VIRGINIA ROAD, P.O. BOX
 9133, CONCORD, MA, 01742-9133
 CLMN Number of Claims: 24
 ECL Exemplary Claim: 1
 DRWN 9 Drawing Page(s)

LN.CNT 2203

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Methods for the production of purified, catalytically active, **recombinant memapsin 2** have been developed-, The substrate and subsite specificity of the catalytically active enzyme have been determined by a method which determines the initial hydrolysis rate of the substrates by using MALDI-TOF/MS. Alternatively, the subsite specificity of memapsin can be determined by probing a library of **inhibitors** with **memapsin 2** and subsequently detecting the bound **memapsin 2** with an antibody raised to **memapsin 2** and an alkaline phosphatase conjugated secondary antibody. The substrate and subsite specificity information was used to design substrate analogs of the natural **memapsin 2** substrate that can **inhibit** the function of **memapsin 2**. The substrate analogs are based on peptide sequences, shown to be related to the natural peptide substrates for **memapsin 2**. The substrate analogs contain at least one analog of an amide bond which is not capable of being cleaved by **memapsin 2**. Processes for the synthesis of substrate analogues including isosteres at the sites of the critical amino acid residues were developed and the more than seventy substrate analogues were synthesized, among which MMI-005, MMI-012, MMI-017, MMI-018, MMI-025, MMI-026, MMI-037, MMI-039, MMI-040, MMI-066, MMI-070, and MMI-071 have **inhibition** constants in the range of 1.4-61.4.times.10.sup.-9 M against **recombinant pro-memapsin 2**. These **inhibitors** are useful in **diagnostics** and for the **treatment** and/or **prevention** of **Alzheimer's** disease.

L6 ANSWER 7 OF 22 USPATFULL on STN

AN 2003:96167 USPATFULL

TI Catalytically active **recombinant** memapsin and methods of use thereof

IN Tang, Jordan J. N., Edmond, OK, United States

Lin, Xinli, Edmond, OK, United States

Koelsch, Gerald, Oklahoma City, OK, United States

Hong, Lin, Oklahoma City, OK, United States

PA Oklahoma Medical Research Foundation, Oklahoma City, OK, United States (U.S. corporation)

PI US 6545127 B1 20030408

AI US 2000-604608 20000627 (9)

PRAI US 1999-141363P 19990628 (60)

US 1999-168060P 19991130 (60)

US 2000-177836P 20000125 (60)

US 2000-178368P 20000127 (60)

US 2000-210292P 20000608 (60)

DT Utility

FS GRANTED

EXNAM Primary Examiner: Borin, Michael; Assistant Examiner: Zhou, Shuba

LREP Hamilton, Brook, Smith & Reynolds, P.C.

CLMN Number of Claims: 18

ECL Exemplary Claim: 7

DRWN 21 Drawing Figure(s); 12 Drawing Page(s)

LN.CNT 2563

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Methods for the production of purified, catalytically active, **recombinant memapsin 2** have been developed. The substrate and subsite specificity of the catalytically active enzyme have been determined. The substrate and subsite specificity information was used to design substrate analogs of the natural **memapsin 2** substrate that can **inhibit** the function of **memapsin 2**. The substrate analogs are based on peptide sequences, shown to be related to the natural peptide substrates for **memapsin 2**. The substrate analogs contain at least one analog of an amide bond which is not capable of being cleaved by

memapsin 2. Processes for the synthesis of two substrate analogues including isosteres at the sites of the critical amino acid residues were developed and the substrate analogues, OMR99-1 and OM99-2, were synthesized. OM99-2 is based on an octapeptide Glu-Val-Asn-Leu-Ala-Ala-Glu-Phe (SEQ ID NO:28) with the Leu-Ala peptide bond substituted by a transition-state **isostere** hydroxyethylene group (FIG. 1). The **inhibition** constant of OM99-2 is 1.6.times.10.sup.-9M against **recombinant pro-memapsin 2**. Crystallography of **memapsin 2** bound to this **inhibitor** was used to determine the three dimensional structure of the protein, as well as the importance of the various residues in binding. This information can be used by those skilled in the art to design new **inhibitors**, using commercially available **software programs** and techniques familiar to those in organic chemistry and enzymology, to design new **inhibitors** to **memapsin 2**, useful in **diagnostics** and for the **treatment** and/or **prevention** of **Alzheimer's** disease.

L6 ANSWER 8 OF 22 USPATFULL on STN
 AN 2003:79303 USPATFULL
 TI 12 human secreted proteins
 IN Ni, Jian, Germantown, MD, UNITED STATES
 Young, Paul E., Gaithersburg, MD, UNITED STATES
 Kenny, Joseph J., Damascus, MD, UNITED STATES
 Olsen, Henrik S., Gaithersburg, MD, UNITED STATES
 Moore, Paul A., Germantown, MD, UNITED STATES
 Wei, Ying-Fei, Berkeley, CA, UNITED STATES
 Greene, John M., Gaithersburg, MD, UNITED STATES
 Ruben, Steven M., Olney, MD, UNITED STATES
 Liu, Ding, Gaithersburg, MD, UNITED STATES
 Crocker, Paul R., Dundee, UNITED KINGDOM
 PI US 2003055231 A1 20030320
 AI US 2001-984130 A1 20011029 (9)
 RLI Continuation-in-part of Ser. No. US 2001-836353, filed on 18 Apr 2001,
 PENDING Continuation-in-part of Ser. No. WO 1999-US25031, filed on 27
 Oct 1999, UNKNOWN
 PRAI US 2000-243792P 20001030 (60)
 US 2000-198407P 20000419 (60)
 US 1998-105971P 19981028 (60)
 DT Utility
 FS APPLICATION
 LREP HUMAN GENOME SCIENCES INC, 9410 KEY WEST AVENUE, ROCKVILLE, MD, 20850
 CLMN Number of Claims: 23
 ECL Exemplary Claim: 1
 DRWN 67 Drawing Page(s)
 LN.CNT 31982
 CAS INDEXING IS AVAILABLE FOR THIS PATENT.
 AB The present invention relates to 12 novel human secreted proteins and
 isolated nucleic acids containing the coding regions of the genes
 encoding such proteins. Also provided are vectors, host cells,
 antibodies, and **recombinant** methods for producing human
 secreted proteins. The invention further relates to **diagnostic**
 and **therapeutic** methods useful for **diagnosing** and
treating disorders related to these novel human secreted
 proteins.

L6 ANSWER 9 OF 22 USPATFULL on STN
 AN 2003:4068 USPATFULL
 TI Method of **preventing** cell death using segments of neural
 thread proteins
 IN Averbach, Paul A., Beaconsfield, CANADA
 PI US 2003004107 A1 20030102
 AI US 2002-146130 A1 20020516 (10)
 PRAI US 2001-290971P 20010516 (60)

DT Utility
FS APPLICATION
LREP HUNTON & WILLIAMS, INTELLECTUAL PROPERTY DEPARTMENT, 1900 K STREET,
N.W., SUITE 1200, WASHINGTON, DC, 20006-1109
CLMN Number of Claims: 14
ECL Exemplary Claim: 1
DRWN 9 Drawing Page(s)
LN.CNT 1698

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Disclosed is a method of **preventing, inhibiting,**
and/or ameliorating cell death and/or tissue necrosis in live tissue by
contacting live tissue with at least a segment of NTP, or homologue,
variant, derivative or mimetic thereof, where the segment of NTP, or
homologue, variant, derivative or mimetic thereof is present in an
amount effective to **prevent, inhibit,** and/or
ameliorate cell death and/or tissue necrosis. The method is capable of
treating conditions requiring **prevention,**
inhibition, and/or amelioration of cell death and/or tissue
necrosis.

L6 ANSWER 10 OF 22 USPATFULL on STN

AN 2003:3410 USPATFULL

TI Method of **preventing** cell death using antibodies to neural
thread proteins

IN Averbach, Paul A., Quebec, CANADA

PI US 2003003445 A1 20030102

AI US 2002-138516 A1 20020506 (10)

PRAI US 2001-288463P 20010504 (60)

DT Utility

FS APPLICATION

LREP HUNTON & WILLIAMS, INTELLECTUAL PROPERTY DEPARTMENT, 1900 K STREET,
N.W., SUITE 1200, WASHINGTON, DC, 20006-1109

CLMN Number of Claims: 12

ECL Exemplary Claim: 1

DRWN 10 Drawing Page(s)

LN.CNT 1705

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Disclosed is a method of **preventing, inhibiting,**
and/or ameliorating cell death and/or tissue necrosis in live tissue
containing neural thread proteins (NTP) by contacting the live tissue
with at least an antibody, antibody fragment or antibody derivative that
recognizes or binds to NTP, where the antibody, antibody fragment or
antibody derivative is present in an amount effective to **prevent**
, **inhibit,** reduce, control and/or ameliorate cell death and/or
tissue necrosis. The method is capable of **treating** conditions
requiring **prevention, inhibition,** reduction, control
and/or amelioration of cell death and/or tissue necrosis caused by the
presence of NTP.

L6 ANSWER 11 OF 22 USPATFULL on STN

AN 2002:294717 USPATFULL

TI Catalytically active **recombinant** memapsin and methods of use
thereof

IN Lin, Xinli, Edmond, OK, UNITED STATES

Koelsch, Gerald, Oklahoma City, OK, UNITED STATES

Tang, Jordan J.N., Edmond, OK, UNITED STATES

PA Oklahoma Medical Research Foundation

PI US 2002164760 A1 20021107

AI US 2001-795903 A1 20010228 (9)

RLI Division of Ser. No. US 2000-604608, filed on 27 Jun 2000, PENDING

PRAI US 1999-141363P 19990628 (60)

US 1999-168060P 19991130 (60)

US 2000-177836P 20000125 (60)

US 2000-178368P 20000127 (60)

US 2000-210292P 20000608 (60)

DT Utility
 FS APPLICATION
 LREP PATREA L. PABST, HOLLAND & KNIGHT LLP, SUITE 2000, ONE ATLANTIC CENTER,
 1201 WEST PEACHTREE STREET, N.E., ATLANTA, GA, 30309-3400
 CLMN Number of Claims: 33
 ECL Exemplary Claim: 1
 DRWN 12 Drawing Page(s)
 LN.CNT 2440
 CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Methods for the production of purified, catalytically active,
recombinant memapsin 2 have been developed.
 The substrate and subsite specificity of the catalytically active enzyme
 have been determined. The substrate and subsite specificity information
 was used to design substrate analogs of the natural **memapsin**
2 substrate that can **inhibit** the function of
memapsin 2. The substrate analogs are based on peptide
 sequences, shown to be related to the natural peptide substrates for
memapsin 2. The substrate analogs contain at least one
 analog of an amide bond which is not capable of being cleaved by
memapsin 2. Processes for the synthesis of two
 substrate analogues including isosteres at the sites of the critical
 amino acid residues were developed and the substrate analogues, OMR99-1
 and OM99-2, were synthesized. OM99-2 is based on an octapeptide
 Glu-Val-Asn-Leu-Ala-Ala-Glu-Phe (SEQ ID NO:28) with the Leu-Ala peptide
 bond substituted by a transition-state **isostere**
 hydroxyethylene group (FIG. 1). The **inhibition** constant of
 OM99-2 is 1.6.times.10.sup.-9 M against **recombinant pro-**
memapsin 2. Crystallography of **memapsin**
2 bound to this **inhibitor** was used to determine the
 three dimensional structure of the protein, as well as the importance of
 the various residues in binding. This information can be used by those
 skilled in the art to design new **inhibitors**, using
 commercially available **software programs** and
 techniques familiar to those in organic chemistry and enzymology, to
 design new **inhibitors** to **memapsin 2**,
 useful in **diagnostics** and for the **treatment** and/or
prevention of Alzheimer's disease.

L6 ANSWER 12 OF 22 USPATFULL on STN
 AN 2002:272761 USPATFULL
 TI Directed evolution of novel binding proteins
 IN Ladner, Robert Charles, Ijamsville, MD, UNITED STATES
 Guterman, Sonia Kosow, Belmont, MA, UNITED STATES
 Roberts, Bruce Lindsay, Milford, MA, UNITED STATES
 Markland, William, Milford, MA, UNITED STATES
 Ley, Arthur Charles, Newton, MA, UNITED STATES
 Kent, Rachel Baribault, Boxborough, MA, UNITED STATES
 PI US 2002150881 A1 20021017
 AI US 2001-781988 A1 20010214 (9)
 RLI Continuation of Ser. No. US 1998-192067, filed on 16 Nov 1998, ABANDONED
 Continuation of Ser. No. US 1995-415922, filed on 3 Apr 1995, PATENTED
 Continuation of Ser. No. US 1993-9319, filed on 26 Jan 1993, PATENTED
 Division of Ser. No. US 1991-664989, filed on 1 Mar 1991, PATENTED
 Continuation-in-part of Ser. No. US 1990-487063, filed on 2 Mar 1990,
 ABANDONED Continuation-in-part of Ser. No. US 1988-240160, filed on 2
 Sep 1988, ABANDONED
 PRAI WO 1989-US3731 19890901
 DT Utility
 FS APPLICATION
 LREP BROWDY AND NEIMARK, P.L.L.C., 624 Ninth Street, N.W., Washington, DC,
 20001
 CLMN Number of Claims: 18
 ECL Exemplary Claim: 1
 DRWN 16 Drawing Page(s)
 LN.CNT 15696

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB In order to obtain a novel binding protein against a chosen target, DNA molecules, each encoding a protein comprising one of a family of similar potential binding domains and a structural signal calling for the display of the protein on the outer surface of a chosen bacterial cell, bacterial spore or phage (genetic package) are introduced into a genetic package. The protein is expressed and the potential binding domain is displayed on the outer surface of the package. The cells or viruses bearing the binding domains which recognize the target molecule are isolated and amplified. The successful binding domains are then characterized. One or more of these successful binding domains is used as a model for the design of a new family of potential binding domains, and the process is repeated until a novel binding domain having a desired affinity for the target molecule is obtained. In one embodiment, the first family of potential binding domains is related to bovine pancreatic trypsin **inhibitor**, the genetic package is M13 phage, and the protein includes the outer surface transport signal of the M13 gene III protein.

L6 ANSWER 13 OF 22 USPATFULL on STN

AN 2002:265848 USPATFULL

TI Biopolymer sequence comparison

IN Toll, Lawrence R., Redwood City, CA, UNITED STATES
Lincoln, Patrick Denis, Woodside, CA, UNITED STATES
Karp, Peter, San Mateo, CA, UNITED STATES
Sonmez, Kemal, Menlo Park, CA, UNITED STATES

PI US 2002146724 A1 20021010

AI US 2001-6492 A1 20011203 (10)

PRAI US 2000-250743P 20001201 (60)

DT Utility

FS APPLICATION

LREP DAVID L. FEIGENBAUM, Fish & Richardson P.C., 225 Franklin Street,
Boston, MA, 02110-2804

CLMN Number of Claims: 71

ECL Exemplary Claim: 1

DRWN 7 Drawing Page(s)

LN.CNT 1796

AB Disclosed are methods, software, and systems for comparing biopolymer sequences. The model includes at least two different characterizations of states of matching between segments of sequences at defined positions. Examples of states of matching include: similarity and dissimilarity between objects, as well as similarity to a reference, e.g., a reference sequence or a sequence profile. A topology of particular match states can be used to identify classes of sequences, e.g., preprohormone sequences.

L6 ANSWER 14 OF 22 USPATFULL on STN

AN 2002:214213 USPATFULL

TI **Inhibitors of memapsin 2** and use thereof

IN Koelsch, Gerald, Oklahoma City, OK, UNITED STATES
Tang, Jordan J.N., Edmond, OK, UNITED STATES
Hong, Lin, Oklahoma City, OK, UNITED STATES
Ghosh, Arun K., River Forest, IL, UNITED STATES

PA Oklahoma Medical Research Foundation (U.S. corporation)

PI US 2002115600 A1 20020822

AI US 2001-845226 A1 20010430 (9)

RLI Division of Ser. No. US 2000-603713, filed on 27 Jun 2000, PENDING

PRAI US 1999-141363P 19990628 (60)

US 1999-168060P 19991130 (60)

US 2000-177836P 20000125 (60)

US 2000-178368P 20000127 (60)

US 2000-210292P 20000608 (60)

DT Utility

FS APPLICATION

LREP Patrea L. Pabst, Arnall Golden & Gregory, LLP, 2800 One Atlantic Center,

1201 West Peachtree Street, Atlanta, GA, 30309-3450

CLMN Number of Claims: 23

ECL Exemplary Claim: 1

DRWN 12 Drawing Page(s)

LN.CNT 2377

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Methods for the production of purified, catalytically active, **recombinant memapsin 2** have been developed. The substrate and subsite specificity of the catalytically active enzyme have been determined. The substrate and subsite specificity information was used to design substrate analogs of the natural **memapsin 2** substrate that can **inhibit** the function of **memapsin 2**. The substrate analogs are based on peptide sequences, shown to be related to the natural peptide substrates for **memapsin 2**. The substrate analogs contain at least one analog of an amide bond which is not capable of being cleaved by **memapsin 2**. Processes for the synthesis of two substrate analogues including isosteres at the sites of the critical amino acid residues were developed and the substrate analogues, OMR99-1 and OM99-2, were synthesized. OM99-2 is based on an octapeptide Glu-Val-Asn-Leu-Ala-Ala-Glu-Phe (SEQ ID NO:28) with the Leu-Ala peptide bond substituted by a transition-state **isostere** hydroxyethylene group (FIG. 1). The **inhibition** constant of OM99-2 is 1.6.times.10.sup.-9 M against **recombinant pro-memapsin 2**. Crystallography of **memapsin 2** bound to this **inhibitor** was used to determine the three dimensional structure of the protein, as well as the importance of the various residues in binding. This information can be used by those skilled in the art to design new **inhibitors**, using commercially available **software programs** and techniques familiar to those in organic chemistry and enzymology, to design new **inhibitors** to **memapsin 2**, useful in **diagnostics** and for the **treatment** and/or **prevention** of Alzheimer's disease.

L6 ANSWER 15 OF 22 USPATFULL on STN

AN 2002:191539 USPATFULL

TI Full-length human cDNAs encoding potentially secreted proteins

IN Milne Edwards, Jean-Baptiste Dumas, Paris, FRANCE

Bougueleret, Lydie, Petit Lancy, SWITZERLAND

Jobert, Severin, Paris, FRANCE

PI US 2002102604 A1 20020801

AI US 2000-731872 A1 20001207 (9)

PRAI US 1999-169629P 19991208 (60)

US 2000-187470P 20000306 (60)

DT Utility

FS APPLICATION

LREP John Lucas, Ph.D., J.D., Genset Corporation, 10665 Sorento Valley Road, San Diego, CA, 92121-1609

CLMN Number of Claims: 29

ECL Exemplary Claim: 1

DRWN 5 Drawing Page(s)

LN.CNT 28061

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The invention concerns GENSET polynucleotides and polypeptides. Such GENSET products may be used as reagents in forensic analyses, as chromosome markers, as tissue/cell/organelle-specific markers, in the production of expression vectors. In addition, they may be used in screening and **diagnosis** assays for abnormal GENSET expression and/or biological activity and for screening compounds that may be used in the **treatment** of GENSET-related disorders.

L6 ANSWER 16 OF 22 USPATFULL on STN

AN 2002:92777 USPATFULL

TI Catalytically active **recombinant** memapsin and methods of use

thereof

IN Tang, Jordan J. N., Edmond, OK, UNITED STATES
 Lin, Xinli, Edmond, OK, UNITED STATES
 Koelsch, Gerald, Oklahoma City, OK, UNITED STATES
 Hong, Lin, Oklahoma City, OK, UNITED STATES

PI US 2002049303 A1 20020425

AI US 2001-796264 A1 20010228 (9)

RLI Division of Ser. No. US 2000-604608, filed on 27 Jun 2000, PENDING

PRAI US 1999-141363P 19990628 (60)
 US 1999-168060P 19991130 (60)
 US 2000-177836P 20000125 (60)
 US 2000-178368P 20000127 (60)

DT Utility

FS APPLICATION

LREP PATREA L. PABST, HOLLAND & KNIGHT LLP, SUITE 2000, ONE ATLANTIC CENTER,
 1201 WEST PEACHTREE STREET, N.E., ATLANTA, GA, 30309-3400

CLMN Number of Claims: 33

ECL Exemplary Claim: 1

DRWN 12 Drawing Page(s)

LN.CNT 2441

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Methods for the production of purified, catalytically active,
recombinant memapsin 2 have been developed.
 The substrate and subsite specificity of the catalytically active enzyme
 have been determined. The substrate and subsite specificity information
 was used to design substrate analogs of the natural **memapsin**
2 substrate that can **inhibit** the function of
memapsin 2. The substrate analogs are based on peptide
 sequences, shown to be related to the natural peptide substrates for
memapsin 2. The substrate analogs contain at least one
 analog of an amide bond which is not capable of being cleaved by
memapsin 2. Processes for the synthesis of two
 substrate analogs including isosteres at the sites of the critical amino
 acid residues were developed and the substrate analogs, OMR99-1 and
 OMR99-2, were synthesized. OMR99-2 is based on an octapeptide
 Glu-Val-Asn-Leu-Ala-Glu-Phe (SEQ ID NO:28) with the Leu-Ala peptide
 bond substituted by a transition-state **isostere**
 hydroxyethylene group (FIG. 1). The **inhibition** constant of
 OMR99-2 is 1.6.times.10.sup.-9 M against **recombinant pro-**
memapsin 2. Crystallography of **memapsin**
2 bound to this **inhibitor** was used to determine the
 three dimensional structure of the protein, as well as the importance of
 the various residues in binding. This information can be used by those
 skilled in the art to design new **inhibitors**, using
 commercially available **software programs** and
 techniques familiar to those in organic chemistry and enzymology, to
 design new **inhibitors** to **memapsin 2**,
 useful in **diagnostics** and for the **treatment** and/or
prevention of **Alzheimer's** disease.

L6 ANSWER 17 OF 22 USPATFULL on STN

AN 2000:7289 USPATFULL

TI Human nucleic acid binding protein

IN Bandman, Olga, Mountain View, CA, United States
 Au-Young, Janice, Berkeley, CA, United States
 Hawkins, Phillip R., Mountain View, CA, United States
 Hillman, Jennifer L., San Jose, CA, United States

PA Incyte Pharmaceuticals, Inc., Palo Alto, CA, United States (U.S.
 corporation)

PI US 6015788 20000118

AI US 1998-195855 19981119 (9)

RLI Division of Ser. No. US 1996-698407, filed on 15 Aug 1996, now patented,
 Pat. No. US 5856128

DT Utility

FS Granted

EXNAM Primary Examiner: Sisson, Bradley; Assistant Examiner: Longton, Enrique D.
LREP Sather, Susan K. Incyte Pharmaceuticals, Inc.
CLMN Number of Claims: 2
ECL Exemplary Claim: 1
DRWN 11 Drawing Figure(s); 11 Drawing Page(s)
LN.CNT 1830

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The present invention provides polynucleotides which identify and encode a novel human nucleic acid binding protein (NABP). The invention provides for genetically engineered expression vectors and host cells comprising the nucleic acid sequences encoding NABP. The invention also provides for the use of substantially purified NABP or its antagonists, in pharmaceutical compositions for the **treatment** of diseases associated with the expression of NABP. Additionally, the invention provides for the use of antisense molecules to NABP in pharmaceutical compositions for **treatment** of diseases associated with the expression of NABP. The invention also describes **diagnostic** assays which utilize **diagnostic** compositions comprising the polynucleotide, fragments or the complement thereof, which hybridize with the genomic sequence or the transcript of polynucleotides encoding NABP or anti-NABP antibodies which specifically bind to NABP.

L6 ANSWER 18 OF 22 USPATFULL on STN

AN 1999:1467 USPATFULL

TI Human nucleic acid binding protein

IN Bandman, Olga, Mountain View, CA, United States

Au-Young, Janice, Berkeley, CA, United States

Hawkins, Phillip R., Mountain View, CA, United States

Hillman, Jennifer L., San Jose, CA, United States

PA Incyte Pharmaceuticals, Inc., Palo Alto, CA, United States (U.S. corporation)

PI US 5856128 19990105

AI US 1996-698407 19960815 (8)

DT Utility

FS Granted

EXNAM Primary Examiner: Wax, Robert A.; Assistant Examiner: Longton, Enrique D.

LREP Billings, Lucy J. Incyte Pharmaceuticals, Inc.

CLMN Number of Claims: 6

ECL Exemplary Claim: 1

DRWN 7 Drawing Figure(s); 7 Drawing Page(s)

LN.CNT 1776

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The present invention provides polynucleotides which identify and encode a novel human nucleic acid binding protein (NABP). The invention provides for genetically engineered expression vectors and host cells comprising the nucleic acid sequences encoding NABP. The invention also provides for the use of substantially purified NABP or its antagonists, in pharmaceutical compositions for the **treatment** of diseases associated with the expression of NABP. Additionally, the invention provides for the use of antisense molecules to NABP in pharmaceutical compositions for **treatment** of diseases associated with the expression of NABP. The invention also describes **diagnostic** assays which utilize **diagnostic** compositions comprising the polynucleotide, fragments or the complement thereof, which hybridize with the genomic sequence or the transcript of polynucleotides encoding NABP or anti-NABP antibodies which specifically bind to NABP.

L6 ANSWER 19 OF 22 USPATFULL on STN

AN 1998:143904 USPATFULL

TI Directed evolution of novel binding proteins

IN Ladner, Robert Charles, Ijamsville, MD, United States

Gutterman, Sonia Kosow, Belmont, MA, United States

Roberts, Bruce Lindsay, Milford, MA, United States

Markland, William, Milford, MA, United States
 Ley, Arthur Charles, Newton, MA, United States
 Kent, Rachel Baribault, Boxborough, MA, United States
 PA Dyax, Corp., Cambridge, MA, United States (U.S. corporation)
 PI US 5837500 19981117
 AI US 1995-415922 19950403 (8)
 RLI Continuation of Ser. No. US 1993-9319, filed on 26 Jan 1993, now
 patented, Pat. No. US 5403484 which is a division of Ser. No. US
 1991-664989, filed on 1 Mar 1991, now patented, Pat. No. US 5223409
 which is a continuation-in-part of Ser. No. US 1990-487063, filed on 2
 Mar 1990, now abandoned which is a continuation-in-part of Ser. No. US
 1988-240160, filed on 2 Sep 1988, now abandoned
 DT Utility
 FS Granted
 EXNAM Primary Examiner: Ulm, John
 LREP Cooper, Iver P.
 CLMN Number of Claims: 43
 ECL Exemplary Claim: 1
 DRWN 16 Drawing Figure(s); 16 Drawing Page(s)
 LN.CNT 15973
 CAS INDEXING IS AVAILABLE FOR THIS PATENT.
 AB In order to obtain a novel binding protein against a chosen target, DNA
 molecules, each encoding a protein comprising one of a family of similar
 potential binding domains and a structural signal calling for the
 display of the protein on the outer surface of a chosen bacterial cell,
 bacterial spore or phage (genetic package) are introduced into a genetic
 package. The protein is expressed and the potential binding domain is
 displayed on the outer surface of the package. The cells or viruses
 bearing the binding domains which recognize the target molecule are
 isolated and amplified. The successful binding domains are then
 characterized. One or more of these successful binding domains is used
 as a model for the design of a new family of potential binding domains,
 and the process is repeated until a novel binding domain having a
 desired affinity for the target molecule is obtained. In one embodiment,
 the first family of potential binding domains is related to bovine
 pancreatic trypsin *inhibitor*, the genetic package is M13
 phage, and the protein includes the outer surface transport signal of
 the M13 gene III protein.
 L6 ANSWER 20 OF 22 USPATFULL on STN
 AN 96:101466 USPATFULL
 TI Directed evolution of novel binding proteins
 IN Ladner, Robert C., Ijamsville, MD, United States
 Guterman, Sonia K., Belmont, MA, United States
 Roberts, Bruce L., Milford, MA, United States
 Markland, William, Milford, MA, United States
 Ley, Arthur C., Newton, MA, United States
 Kent, Rachel B., Boxborough, MA, United States
 PA Protein Engineering Corporation, Cambridge, MA, United States (U.S.
 corporation)
 PI US 5571698 19961105
 AI US 1993-57667 19930618 (8)
 DCD 20100629
 RLI Continuation of Ser. No. US 1991-664989, filed on 1 Mar 1991, now
 patented, Pat. No. US 5223409 which is a continuation-in-part of Ser.
 No. US 1990-487063, filed on 2 Mar 1990, now abandoned which is a
 continuation-in-part of Ser. No. US 1988-240160, filed on 2 Sep 1988,
 now abandoned
 DT Utility
 FS Granted
 EXNAM Primary Examiner: Ulm, John
 LREP Cooper, Iver P.
 CLMN Number of Claims: 83
 ECL Exemplary Claim: 1
 DRWN 16 Drawing Figure(s); 16 Drawing Page(s)

LN.CNT 15323

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB In order to obtain a novel binding protein against a chosen target, DNA molecules, each encoding a protein comprising one of a family of similar potential binding domains and a structural signal calling for the display of the protein on the outer surface of a chosen bacterial cell, bacterial spore or phage (genetic package) are introduced into a genetic package. The protein is expressed and the potential binding domain is displayed on the outer surface of the package. The cells or viruses bearing the binding domains which recognize the target molecule are isolated and amplified. The successful binding domains are then characterized. One or more of these successful binding domains is used as a model for the design of a new family of potential binding domains, and the process is repeated until a novel binding domain having a desired affinity for the target molecule is obtained. In one embodiment, the first family of potential binding domains is related to bovine pancreatic trypsin **inhibitor**, the genetic package is M13 phage, and the protein includes the outer surface transport signal of the M13 gene III protein.

L6 ANSWER 21 OF 22 USPATFULL on STN

AN 95:29292 USPATFULL

TI Viruses expressing chimeric binding proteins

IN Ladner, Robert C., Ijamsville, MD, United States

Guterman, Sonia K., Belmont, MA, United States

Roberts, Bruce L., Milford, MA, United States

Markland, William, Milford, MA, United States

Ley, Arthur C., Newton, MA, United States

Kent, Rachel B., Boxborough, MA, United States

PA Protein Engineering Corporation, Cambridge, MA, United States (U.S. corporation)

PI US 5403484 19950404

AI US 1993-9319 19930126 (8)

RLI Division of Ser. No. US 1991-664989, filed on 1 Mar 1991, now patented, Pat. No. US 5223409 which is a continuation-in-part of Ser. No. US 1990-487063, filed on 2 Mar 1990, now abandoned which is a continuation-in-part of Ser. No. US 1988-240160, filed on 2 Sep 1988, now abandoned

PRAI WO 1989-3731 19890901

DT Utility

FS Granted

EXNAM Primary Examiner: Hill, Jr., Robert J.; Assistant Examiner: Ulm, John D.

LREP Cooper, Iver P.

CLMN Number of Claims: 49

ECL Exemplary Claim: 1

DRWN 16 Drawing Figure(s); 16 Drawing Page(s)

LN.CNT 14368

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB In order to obtain a novel binding protein against a chosen target, DNA molecules, each encoding a protein comprising one of a family of similar potential binding domains and a structural signal calling for the display of the protein on the outer surface of a chosen bacterial cell, bacterial spore or phage (genetic package) are introduced into a genetic package. The protein is expressed and the potential binding domain is displayed on the outer surface of the package. The cells or viruses bearing the binding domains which recognize the target molecule are isolated and amplified. The successful binding domains are then characterized. One or more of these successful binding domains is used as a model for the design of a new family of potential binding domains, and the process is repeated until a novel binding domain having a desired affinity for the target molecule is obtained. In one embodiment, the first family of potential binding domains is related to bovine pancreatic trypsin **inhibitor**, the genetic package is M13 phage, and the protein includes the outer surface transport signal of the M13 gene III protein.

L6 ANSWER 22 OF 22 USPATFULL on STN
AN 93:52487 USPATFULL
TI Directed evolution of novel binding proteins
IN Ladner, Robert C., Ijamsville, MD, United States
Guterman, Sonia K., Belmont, MA, United States
Roberts, Bruce L., Milford, MA, United States
Markland, William, Milford, MA, United States
Ley, Arthur C., Newton, MA, United States
Kent, Rachel B., Boxborough, MA, United States
PA Protein Engineering Corp., Cambridge, MA, United States (U.S.
corporation)
PI US 5223409 19930629
AI US 1991-664989 19910301 (7)
RLI Continuation-in-part of Ser. No. US 1990-487063, filed on 2 Mar 1990,
now abandoned And a continuation-in-part of Ser. No. US 1988-240160,
filed on 2 Sep 1988, now abandoned
DT Utility
FS Granted
EXNAM Primary Examiner: Hill, Jr., Robert J.; Assistant Examiner: Ulm, John D.
LREP Cooper, Iver P.
CLMN Number of Claims: 66
ECL Exemplary Claim: 1
DRWN 16 Drawing Figure(s); 16 Drawing Page(s)
LN.CNT 15410
CAS INDEXING IS AVAILABLE FOR THIS PATENT.
AB In order to obtain a novel binding protein against a chosen target, DNA
molecules, each encoding a protein comprising one of a family of similar
potential binding domains and a structural signal calling for the
display of the protein on the outer surface of a chosen bacterial cell,
bacterial spore or phage (genetic package) are introduced into a genetic
package. The protein is expressed and the potential binding domain is
displayed on the outer surface of the package. The cells or viruses
bearing the binding domains which recognize the target molecule are
isolated and amplified. The successful binding domains are then
characterized. One or more of these successful binding domains is used
as a model for the design of a new family of potential binding domains,
and the process is repeated until a novel binding domain having a
desired affinity for the target molecule is obtained. In one embodiment,
the first family of potential binding domains is related to bovine
pancreatic trypsin inhibitor, the genetic package is M13
phage, and the protein includes the outer surface transport signal of
the M13 gene III protein.

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---Logging off of STN---

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STN INTERNATIONAL LOGOFF AT 16:46:14 ON 05 AUG 2003